

BKL Annotation details of one docket matching protein

**Human SNAI1** Snail 1 (*Drosophila*) homolog, a zinc-finger transcriptional repressor, represses expression of E-cadherin (CDH1) and aromatase (CYP19), may play a role in carcinoma and melanoma progression by repressing CDH1

**Disease**

**Therapeutic Target:**

decreased expression of SNAI1 mRNA may prevent abnormal cell differentiation associated with Carcinoma [2000](#) ([10655587](#))

**Diagnostic Marker:**

decreased expression of SNAI1 mRNA may correlate with Breast Neoplasms [2001](#) ([11245431](#))  
increased expression of SNAI1 protein correlates with breast ductal carcinoma associated with Breast Neoplasms [2002](#) ([12082640](#))  
increased expression of SNAI1 protein correlates with breast ductal carcinoma [2002](#) ([12082640](#))  
increased expression of SNAI1 mRNA may correlate with increased negative regulation of transcription from Pol II promoter associated with Melanoma [2001](#) ([11323412](#))  
increased expression of SNAI1 mRNA may correlate with malignant form of Melanoma [2001](#) ([11323412](#))

**Negative Correlation:**

SNAI1 gene does not correlate with Craniosynostoses [1999](#) ([10585766](#)) [1999](#) ([10543399](#))

**Phenotype**

**Title line  
phrases**

**Membership:**

contains an N-terminal SNAG domain [2003](#) ([12579345](#))  
member of the SNAG zinc finger protein subfamily of zinc finger proteins [2003](#) ([12579345](#))

**Biological Process/Role:**

represses expression of E-cadherin (CDH1) [2000](#) ([10655587](#))  
represses expression of aromatase (CYP19) [2001](#) ([11245431](#))

**Role in Disease:**

involved in tumor progression [2000](#) ([10655587](#))  
upregulated in melanoma cells [2001](#) ([11323412](#))  
expression inversely correlates with the grade of differentiation of breast carcinoma [2002](#) ([12082640](#))  
downregulated in breast cancer cell lines [2001](#) ([11245431](#))

**Synonyms**

SNA  
SLUGH2  
SNAH  
dJ710H13.1

**Cognate**

Mouse Snail \*

**members**      Rat Snail

**GO**      GO ontology: transcriptional repressor activity *Experimental (E) 2001 (11245431)*  
 GO ontology: specific transcriptional repressor activity *Experimental (E) 2000 (10655587)*  
 GO ontology: specific transcriptional repressor activity *Experimental (E) 2000 (10655586)*  
 GO ontology: DNA binding *Experimental (E) 2001 (11245431)*  
 GO ontology: cartilage condensation *Unspecified Evidence (?) 1992 (1295727)*  
 GO ontology: neurogenesis *Unspecified Evidence (?) 1992 (1295727)*  
 GO ontology: negative regulation of transcription from Pol II promoter *Experimental (E) 2001 (11245431)*  
 GO ontology: negative regulation of transcription from Pol II promoter *Experimental (E) 2000 (10655587)*  
 GO ontology: mesoderm cell fate determination *Unspecified Evidence (?) 1992 (1295727)*  
 GO ontology: mesoderm cell fate determination *Unspecified Evidence (?) 1992 (1483390)*

**Expression**      Body: mammary gland/breast \* Cell types: fibroblasts \* Cell origin: cell line \*  
 Techniques: rt-PCR *Experimental (E) 2001 (11245431)*  
 Body: lung \* developmental stage: adult \* Techniques: Northern analysis  
*Experimental (E) 1999 (10585766)*  
 Body: liver \* developmental stage: adult \* Techniques: Northern analysis  
*Experimental (E) 1999 (10585766)*  
 Body: mammary gland/breast \* Tumors: tumor \* Cell origin: cell line \* Regulation:  
 downregulated \* Techniques: rt-PCR *Experimental (E) 2001 (11245431)*  
 Body: skeletal muscle \* developmental stage: adult \* Techniques: Northern analysis  
*Experimental (E) 1999 (10585766)*  
 Body: placenta \* Techniques: Northern analysis *Experimental (E) 1999 (10585766)*  
 Cell origin: cell line \* Techniques: in situ hybridization \* Tumors: melanoma  
*Experimental (E) 2000 (10655586)*  
 Cell origin: cell line \* Techniques: Northern analysis \* Tumors: tumor *Experimental (E) 2000 (10655587)*  
 Cell origin: cell line \* Cell types: fibroblasts \* Techniques: Northern analysis  
*Experimental (E) 2000 (10655587)*  
 Body: brain \* developmental stage: adult \* Techniques: Northern analysis  
*Experimental (E) 1999 (10585766)*  
 Body: heart \* developmental stage: adult \* Techniques: Northern analysis  
*Experimental (E) 1999 (10585766)*  
 Tumors: carcinoma \* Cell origin: cell line \* Techniques: in situ hybridization \* Body:  
 mammary gland/breast *Experimental (E) 2000 (10655586)*  
 Cell origin: cell line \* Tumors: melanoma \* Techniques: rt-PCR *Experimental (E) 2000 (10655586)*

Tumors: carcinoma \* Cell origin: cell line \* Body: mammary gland/breast \*  
 Techniques: rt-PCR *Experimental (E) 2000* ([10655586](#))  
 Cell types: fibroblasts \* Cell origin: primary cells in culture \* Techniques: rt-PCR \*  
 Body: skin *Experimental (E) 2001* ([11323412](#))  
 Cell types: melanocytes \* Degree: not \* Cell origin: primary cells in culture \*  
 Techniques: rt-PCR *Experimental (E) 2001* ([11323412](#))  
 Cell origin: cell line \* Tumors: melanoma \* Techniques: rt-PCR \* Regulation:  
 upregulated *Experimental (E) 2001* ([11323412](#))  
 Body: kidney \* developmental stage: embryo-fetus \* Techniques: Northern analysis  
*Experimental (E) 1999* ([10543399](#))  
 developmental stage: embryo-fetus \* Techniques: Northern analysis \* Misc. Organ/Cell  
 Type: several tissues *Experimental (E) 1999* ([10543399](#))  
 Body: mammary gland/breast \* Cell types: epithelium/epithelial cells \* Cell origin: cell  
 line \* Techniques: rt-PCR *Experimental (E) 2001* ([11245431](#))

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